## AMENDMENTS TO THE SPECIFICATION

Please amend the specification as follows:

Please amend the paragraph on page 12, lines 24-30:

For example, synthesis of the first cDNA strand may be carried out in the presence of primers having appropriate restriction sites in order to allow a subsequent cloning in the proper orientation with respect to the expression vector promoter. As restriction sites, any known restriction site may be used. As a primer, for instance the following primer, 50 nucleotides long may be used: (SEQ ID NO: 181)

Please amend the paragraph bridging pages 12 and 13:

After two-strand synthesis, the cohesive ends of the double stranded cDNA are filled (blunt end) and the cDNA ends are then ligated using a suitable DNA adaptor sequence. The DNA adaptor sequence should contain a restriction site which should be different from the restriction site used in the primer for the synthesis of the first cDNA strand. The DNA adaptor may comprise for example complementary 9- or 13-mer oligonucleotides, whose ends represent the cohesive end of a restriction site. These ends may be for example a EcoRI-site:

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5' XXXXXGGCACGAG 3' (SEQ ID NO: 182)

3' XCCGTGCTC 5'

Please amend the paragraphs on page 24, lines 3-32:

pKS+ANotI is cleaved with PstI and BamHI and the DNA oligonucleotide, synthesized from the pK3/pK4 primer pair described below, is ligated in the opened plasmid. The pKS+neu plasmid thus prepared contains between PstI and BamHI restriction sites, the following novel restriction sites NotI, StuI, SfiI and NcoI (i.e. PstI-NotI-StuI-SfiI-NcoI-BamHI)

- 5'-GCGGCCGCAAGGCCTCCATGGCCG-3' PK3 (SEQ ID NO: 183)
- 5'-GATCCGGCCATGGAGGCCTTGCGGCCGCTGCA-3' PK4 (SEQ ID NO: 184)

The URA3 gene of S.cerevisiae is amplified via PCR, by use of the primer-pair PK9 and PK10, described below, and an Ycplac33 vector DNA (Gietz, R. D. and Sugino, A. (1988) Gene 74: 527-534) as matrix. The amplified DNA is cleaved with BamHI and NotI and subsequently inserted in pKS+neu which has been cleaved by BamHI and NotI. The plasmid thus obtained is named pPK9/10.

..NotI..
5'-ATCTGCAGCGGCCG<u>CAAACATGAGAATTGGGTAATAACTG</u>-3' PK9

PstI

(SEQ ID

NO:185)

..SfiI..

## 5'-ATGGATCCGGCCATGGAGGCCTTCAAGAATTAGCTTTTCAATTCATC-3'

BamH1

PK10 (SEQ ID

NO: 186)

2) Preparation of the deletion cassette

The 5'-region of ORF YML114c was amplified by PCR using genomic DNA of S.cerevisiae as template and both primers YML114c-Asp718 and YLM114c-EcoRI, described below.

YML114c-Asp718: 5'-GCTGGTACCCGTCGGTCTCTTTACC-3'(SEQ ID NO: 187)

YLM114c-EcoRI: 5'-TTGGAATTCATTGCCCTTTATGAGTCC-3'(SEQ ID NO: 188)

Please amend the paragraph on page 25, lines 1-9:

The 3'region of ORF YML114c was amplified by PCR using genomic DNA of S.cerevisiae as template and both primers YML114c-BamHI and YLM114c-SacI, described below.

YML114c-BamHI:5'-ATCGGATCCGCCAACAATGACAGCG-3'(SEQ ID NO: 189)

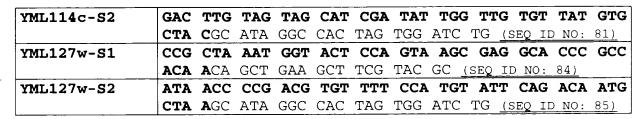
YLM114c-SacI: 5'-GTTGAGCTCTGAGCGTTTGTCCTTG-3'(SEQ ID NO: 190)

The PCR fragment was subsequently cut with BamHI and SacI. The resulting 535bp fragment was inserted in plasmid pYML114c-A linearized with BamHI and SacI generating pYML114c-B.

TABLE 2: Primers used for gene deletions

Gene deletion	ns on chromosome 13
Name	Sequence 5'-3'
YDR472w-S1	ATG TCT CAA AGA ATA ATT CAA CCA AGC GCA TCT GAC
	CAA CCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 167)
YDR472w-S2	AGC CAA ATC TCA AAC CTT CCC TGT CAA GCA CTT GCC
	TGT CGC ATA GGC CAC TAG TGG ATG TG (SEQ ID NO:
	177)
YDR499w-S1	ATG AGA CGA GAA ACG GTG GGT GAA TTT TCT TCA GAT
	GAC GCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 95)
YDR499w-S2	CGT ACT TTA CTT GCA TTA TTC TCC CCG TTC TTT TAT
	TCA AGC ATA GGC CAC TAG TGG ATG TG (SEQ ID NO:
	175)
YMR049c-S1	CAG ACT ATT GAT TAC TTT ATG ACC GGT TAG TTT CTT
	TAG TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 170)
YMR049c-S2	TCT GTT CTA ACA TAA CTA GGT CAA TGA TGG CTA AGA
	ACA AGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
10m124 01	128)
YMR134w-S1	GCA AAG TGT GGT ATA GAA AAA GAA CCA AAG GCC GGT
10m124 00	ATG TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 139)
YMR134w-S2	TGT GTG TGT GCC TAC CTG CAT GTA TGC ATT TAG CAA
	TTG AGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO: 150)
YML023c-S1	CAC GCA ATG GTG CAC ATT ATT TTG TTG AAC TCA CTG
IMIO23C-SI	AGA ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 159)
YML023c-S2	ATT AGT TAC TTA TTC TAT AAT TAC ACT TTT ATC ATG
IMD025C-52	AAC GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO:
	173)
YML049c-S1	AAT TCC TGC TCA TTC AAG GAA AGT CTC AGG AAA TTT
	TCA CCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 106)
YML049c-S2	ACT CCT GCA TCG GAC ACT TCG TCG ATC TGG AAG CAG
	GGT CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO: 19)
YML077w-S1	ATG GGG ATA TAT TCA TTT TGG ATC TTT GAT AGG CAT
	TGT ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 28)
YML077w-S2	TTC TAT TGG TGA TCT TTC TTG TCC CTT GAC CTC TCA
	TTT CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO: 39)
YML093w-S1	GCT AAC TTA AAT ATG GCA AAA AAG AAA TCT AAG AGC
	AGA TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 50)
YML093w-S2	CAA AGG ATC AAT AAC TTG GCC TGG CTT AGT CAT GAT
	TCT CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO: 61)
YML114c-S1	AAC GTG TAA TTG AGG GAC TCA TAA AGG GCA ATG ACT
	TCC ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 71)
	1 - C - C - C - C - C - C - C - C - C -

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Gene deletions	on (	chro	nosoı	me 13	3							
Name	Sequ	ence	5'-	-3'								
YMR032w-S1	CTA	CAG	TTA	TGA	AGC	TTG	TTT	TTG	GGA	CCC	AAA	CGA
	CAA	TCA								D NO:		
YMR032w-S2		AAA									GAT	
	AGA										NO:	
YMR093w-S1	ATG		ACT								TCG	AAG
	<b></b>							=		D NO:		
YMR093w-S2		CAC			AGT						GAT	
	TCT	<b>C</b> GC									O NO:	
YMR131c-S1	CTT	TAA		CCT							ATA	ATT
	GTG									D NO:		
YMR131c-S2	GGT							_	_		GAG	
	1										: ОИ С	
YMR185w-S1	ATC						_		_		CCG	AAG
	CTC	<b>A</b> CA	GCT				TAC			D NO:		
YMR185w-S2	GTA	ATG									AAA	
	<del></del>										: ON C	
YMR212c-S1	CCT	CTT		_		_					TTT	GCG
										D NO:		
YMR212c-S2		ATG									CTG	
	AAT										О ИО:	
YMR213w-S1	ATA										GGA	GGA
		·								D NO:		
YMR213w-S2	GCT		ACT								ATT	
	<del></del>	<b>A</b> GC									NO:	
YMR218c-S1	GAC		AAT								CAA	CTT
	<del>                                     </del>	CCA					TAC			D NO:		
YMR218c-S2		GGC									TAA	
	GCT	<b>T</b> GC	ATΑ	A GG(	C CA	C TA	AG 110	∃G A	TC T	'G <u>(s</u>	EQ ID	NO:
YMR281w-S1	100)	220	222	3 CM	ma a	A III C	220	N III C	mmc	A C C	CGT	7.77
IMKZ01W-D1	İ									D NO:		ACA
YMR281w-S2	<del> </del>										GAA	NGC.
IMRZ61W-5Z	TTG										EQ ID	
	102)	<b>1</b> GC	AIL	1 001	CA		10 10	JO A	10 1	0 15	EQ ID	NO.
YMR288w-S1	<del></del>	AAC	CTG	CAG	AAA	GAA	GCT	GCA	CGT	ATT	GGT	GAG
										D NO:		
YMR288w-S2	<del>                                     </del>										GTG	GTG
	AAG	TGC	ATA	GG(	C CA	C TA	G TO	GG A	TC I	G (s	EQ ID	NO:
	104)											
YMR290c-S1	TGA	GTT	TTA	CGT	CTT	TTG	GTA	TTT	GGC	GTT	TTT	CCA
	CTG	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ I	D NO:	105)	
YMR290c-S2	GAT	AAG	CTG	AGC	AAT	ATT	AAC	AGG	AGA	AGT	ATG	GCT
	ACC	<b>C</b> GC	ATA	GG(	C CA	C TA	AG TO	GG A	TC T	'G <u>(s</u>	EQ ID	NO:
	107)											

Bil

YMR211w-S1	AGA	GAG	CAA	ACC	ATT	TGA	CTA	CTC	: AA	r TCT	TCA	ATA
	TAC	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ :	ID NO:	108)	
YMR211w-S2	ATT	TCA	ATC	ATC	TTA	CTC	CGT	GAA	TC	A GGT	TCG	GAA
	TGA	TGC	ATA	GG	C CA	C TA	G T	GG A	TC '	TG (S	EQ ID	NO:
	109)											

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Gene deletions	on	chror	noso	me 4								
Name	Sequ	ience	5'-	-3'								
YDR196c-S1	ATG	CTT	ATG	ATC	AAA	TTG	TGT	TAT	ACT	TCA	AGG	ACA
	AAA	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	NO:	110)	
YDR196c-S2	TTT	CAA	TCT	GTT	CGT	ATA	AGT	CAA	CCA	ATG	TGC	TGT
	TAT	TGC	ATA	A GG	C CA	.С ТА	AG TO	GG A	TC T	G <u>(s</u> :	EQ ID	NO:
	111)		<u> </u>									
YDR299w-S1											GAT	ATC
	GCC	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	O NO:	112)	
YDR299w-S2	CAA	AGA	TTT	GGA	TAT	CAT	CGT	$\mathbf{T}\mathbf{T}\mathbf{T}$	TAA	CAG	CCT	CTA
	ATT	<b>C</b> GC	ATA	A GG	C CA	.C TA	AG TO	GG A'	TC T	G <u>(s</u>	EQ ID	NO:
	113)											
YDR365c-S1	CTG	GAG	AGA	ACC	CAA	AGA	AGG	AAG	GTG	TAG	ATG	CTA
	GGT	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	ONO:	114)	
YDR365c-S2	TTA	GTA	TGC	TTT	TTA	TTA	ACA	GAT	TTC	AAC	TTG	CTT
	TTC	TGC	ATI	A GG	C CA	.C TA	AG TO	GG A	TC T	G <u>(s</u>	EQ ID	NC):
	115)											
YDR396w-S1	CAG	ATA	CAC	TAT	TGT	GGT	GTA	ATC	TGG	ACC	TTG	ACT
	GTC	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	ONO:	116)	
YDR396w-S2	TAG	AGA	AAA	CAC	TGA	ATG	ATC	TTA	GCG	ACC	GTA	CAA
	AAG	<b>A</b> GC	ATA	A GG	C CA	C TA	AG TO	GG A	TC T	G <u>(s</u>	EQ ID	NO:
	118)											
YDR407c-S1	TTC	TTA	AGC	ATT	TCC	CAA	GCT	ATG	TTG	GCC	CAT	CTA
	AGA	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	O NO:	119)	
YDR407c-S2	AAT	AAC	AGA	CAA	GAT	AAC	GTT	TTC	AGA	GTC	GAA	CTG
	GAT	<b>T</b> GC	ATA	A GG	C CA	C TA	AG TO	GG A	TC T	G <u>(s</u>	EQ ID	NO:
	120)											
YDR416w-S1	ACT	TAC	ATG	GAA	AAG	ATA	TAT	CGA	GTA	TTG	GAA	AGA
	GGA	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	O NO:	121)	
YDR416w-S2	TCA	AAT	ATC	TAG	TTC	TAT	TTC	ATC	TGG	ATT	AAT	CGA
	ATA	<b>T</b> GC	ATA	A GG	C CA	C TA	AG TO	GG A	TC T	G <u>(s</u>	EQ ID	NO:
	122)						~					
YDR449c-S1	CAC	ATC	ACC	GAT	TTC	TAA	TAA	TGT	CGA	AGA	CAA	GAT
	ACT	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	ONO:	123)	
YDR449c-S2	ATA	ATT	AAA	TCT	AGA	ATT	TTA	TAC	CTA	GGA	TCA	TCT
	TCT	<b>G</b> GC	ATA	A GG	C CA	C TA	AG TO	GG A	TC T	G <u>(s</u>	EQ ID	NO:
	124)		.,									
YDR141c-S1	TTC	GTA	ATC	TTT	GAA	TTC	TGC	GAT	TTC	ATC	TAC	CAG
	CGC	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	O NO:	125)	
YDR141c-S2	CAC	TAA	AGC	CCC	TTA	CAA	TTG	ACT	CAA	ATA	ATA	AAC
	AAC	<b>T</b> GC	ATA	A GG	C CA	C TA	AG TO	GG A	TC T	G <u>(s</u>	EQ ID	NO:
	126)											
YDR324c-S1	AAG	AAG	CCT	GAA	AAT	ACG	AAA	CAA	ACC	GGT	GAA	GAT
	GAC	<b>C</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	O NO:	127)	
YDR324c-S2	AAA	CAC'	raa	CTT	TGG	TTG	AAT	AAA	CGC	CTT	TTG	TTT
	GGA	<b>G</b> GC	ATA	A GG	C CA	C TA	AG TO	GG A	TC T	G <u>(s</u>	EQ ID	NO:
	129)											

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YDR325w-S1	GAC	ATT	AAT	ACG	AAA	ATC	TTT	AAC	TCA	GTT	GCT	GAA
	GTA	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u> :	SEQ II	NO:	130)	
YDR325w-S2	ACC	TCG	CTG	AAA	GAC	TCT	GAA	TCC	TTA	TCT	TCT	TCA
	TCT	<b>A</b> GC	ATA	GG(	C CA	C TA	G TO	GG A'	rc T	G <u>(S</u>	EQ ID	NO:
	131)											
YDR398w-S1	ATG	GAT	TCT	CCT	GTT	CTA	CAG	TCC	GCT	TAT	GAC	CCA
	TCA	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	132)	
YDR398w-S2	AAC	GTC	ACT	ATA	TCC	GGC	TTC	CTC	CTC	GCC	GTC	GCT
	CTG	<b>C</b> GC	ATP	GG(	C CA	C TA	G TO	G A'	rc r	G <u>(SI</u>	EQ ID	NO:
	133)											

BL

Gene deletio	ns on chromosome 4
Name	Sequence 5'-3'
YDR246w-S1	ATG GCC ATC GAA ACA ATA CTT GTA ATA AAC AAA TC
	GGC GCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 134)
YDR246w-S2	AAC AGG TTA GAT CTT ATA GGC ATT TCC ATT GAG TA
	GAT GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	135)
YDR236c-S1	CTA AAA TAT TGA ACT TGA CCC TGG CCC CAT AAA AA
	CAT TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 136)
YDR236c-S2	TTG AAG TGT TGA TGT TTACGT GGA CTA TTT ATG TT
	CGT TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	137)
YDR361c-S2	TTA CCA AGT GGA AAT TTC TGT TTC CAA TTC ATC GA
	ACT TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	138)
YDR361c-S1	GGT TCA AGC TAT CAA ATT AAA TGA TTT AAA AAA TA
IIDD 2 67 61	GAA GCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 140)
YDR367w-S1	ATC TGC GTA CTT TAT ACA ATC GAT ACC ATT TCC AC
	TGT TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 141)
YDR367w-S2	GTT TTG TTC TAC GTC ATC CCT ATC AAC TAA ATA TT
	GGG GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
YDR339c-S1	TAT GGG TAA AGC TAA GAA AAC AAG AAA GTT TGG CC
IDR339C-SI	CGT ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 143)
YDR339c-S2	TAA AAG ACA TCT GGC AAT TTT TCA ATG ACG TAT GC
IDK359C-52	TGA CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	144)
YDR413c-S1	TTC TTT GGT TTA TTC TTC GTT CAT TTT TGG TCA AA
	ATC TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 145)
YDR413c-S2	ACA AAA GAA AGC ACA AGA GTT TAT TAA GGA GCA GG
IDKIIIO DI	AAG GGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	146)
YDR429c-S1	TCT AGA TCT ATC ATT ACA TAC AAG ATT GAA GAC GG
	GTC ACA GCT GAA GCT TCG TAC GC (SEQ ID NO: 147)
YDR429c-S2	TTT CTT TGT TTC TAA CGA CAG AAA CTC TTG GAA TG
	GTG CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	148)
YDR468c-S1	GTC ACA ATA CTG CTG GTG ATG ACG ATC AAG AGG AG
_	AAA TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 149)
YDR468c-S2	CAA GAC GAC AAT AAG AAG TCC TAT ACA ACA ATC GT
	GTA TGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	151)
YDR489w-S1	ACT ACC CAC AGA GAT GCA AAT ACA ATA GTG GGT TC
	TCC TCA GCT GAA GCT TCG TAC GC (SEQ ID NO: 152)
YDR489w-S2	AGT CGG GCT CAT CTA TCA TGT TTA CGC TAC CTT CT
	TAT CGC ATA GGC CAC TAG TGG ATC TG (SEQ ID NO
	178)

B6 cons

YDR527w-S1	ATG	GAC	TTA	CTG	GGC	GAT	ATA	GTG	GAG	AAA	GAT	ACA
	TCT	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u> :	SEQ II	NO:	<u>179)</u>	
YDR527w-S2	CCC	CAC	CGC	CTT	GTT	TCC	ATA	ACC	AAA	GTG	CAT	CAA
	TAG	<b>C</b> GC	ATA	GG(	C CA	C TA	G TO	GG A	rc To	G <u>(S</u> I	EQ ID	NO:
	<u>153)</u>			_								
YDR288w-S1	ATG	AGT	TCT	ATA	GAT	AAT	GAC	AGC	GAT	GTG	GAT	TTA
	ACA	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u> :	SEQ II	NO:	154)	
YDR288w-S2	GCC	CAT	GAT	TTC	TTG	CAC	CAA	TTT	TTC	AAG	AGA	CTC
	TAG	TGC	ATA	GG(	C CA	C TA	AG TO	G A	TC TO	G <u>(si</u>	EQ ID	NO:
	155)											

B b

Gene deletions	on (	chro	nosor	ne 4								
Name	Sequ	ence	5'-	٠3 '								
YDR201w-S1	CCC	ATG	тст	GGA	CTA	TTC	AGA	GCA	TCA	TCG	TCA	TCC
	ATA	<b>C</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	NO:	156)	
YDR201w-S2	AAA	AGG	GTT	TTC	CGT	TTA	GTT	CCC	GAA	TAT	GAT	GTT
	GAA	<b>A</b> GC	ATA	GG(	C CA	$C$ $T^{\mu}$	G TO	GG A	TC TO	G <u>(s</u> :	EQ ID	NO:
	157)											
YDR434w-S1	ATG	TCC	AAT	GCA	AAT	CTA	AGA	AAA	TGG	GTT	GGT	TTT
	TGC	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u> :	SEQ II	NO:	158)	
YDR434w-S2	TAA	AGG	TAA	ATA	CAC	AGC	TAT	CAT	GTG	CTC	TTG	TGG
	GAA	<b>G</b> GC	ATA	GG	C CA	C TA	AG TO	GG A	rc To	G <u>(s</u> e	O ID	NO:
	160)						_					
YDR181c-S1	AGG	ATA	AAC	CCA	AAT	GCT	GGA	CAT	CTA	AGG	AAA	TCT
	AAG	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u> :	SEQ II	NO:	161)	
YDR181c-S2	TAG	TTG	GGT	TTG	AAT	CGT	TAT	CAC	GGG	AGA	ACA	TTG
	CTT	TGC	ATA	GG(	C CA	C $TP$	AG TO	GG A'	TC T	G <u>(s</u>	EQ ID	NO:
	162)											
YDR531w-S1	ATG	CCG	CGA	ATT	ACT	CAA	GAG	ATA	TCT	TAC	AAT	TGC
	GAT	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u> :	SEQ II	NO:	163)	
YDR531w-S2	AAA	TAA	GCT	ATT	TGC	CCA	ATA	TTG	TTG	GAG	ATG	GCG
	AAT	A <b>GC</b>	ATA	GG	C CA	C TA	AG T	GG A	rc T	G <u>(s</u> :	EQ ID	NO:
	164)											

B6 Const

Gene deletions	on	chro	noson	me 1	2							
Name	Sequ	ence	5'-	-3'								
YLR186w-S1	CTA	GTC	ACC	AAG	AAG	AAA	ACC	CGT	AAA	ATC	GTA	GGT
	CAT	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	ONO:	165)	
YLR186w-S2	ATA	CAA	AGA	GGA	TGC	CAA	GTA	GAC	TTA	AAC	ACT	ATA
	AAA	TGC	ATI	A GG	C CA	C TP	G T	GG A	TC T	G <u>(s</u> :	EQ ID	NO:
	166)											
YLR215c-S1								_		_	AAC	TTT
									SEQ II			
YLR215c-S2	AGC										GCT	
	ACG	<b>A</b> GC	ATF	A GG	C CA	C TP	G TO	GG A	TC T	G <u>(s</u> :	EQ ID	<u>NO:</u>
	169)											
YLR222c-S1	CTC										AAT	ATG
	+								SEQ II			
YLR222c-S2										_	ACC	
		<b>A</b> GC	ATA	A GG	C CA	C TP	AG TO	GG A	TC T	G <u>(s</u> :	EQ ID	NO:
TT DO 42 01	174)	mam		amm	COM		3.000	OM.	mm a	003	COM	001
YLR243w-S1		_		_							CCT	GCA
TT DO 42 GO	GGT					-			SEQ II			
YLR243w-S2					_		_				TTA	
		AGC	ATA	A GG	C CA	.C 17	AG T	JG A	TC T	G (S.	EQ ID	NO:
YLR272c-S1	171)	CCC	TOT	CCC	እርጥ	ጥጥር	TCA	CTC	ጥጥር	CAA	CTA	<b>አ</b> ጥጥ
ILRZ/ZC-SI									SEQ II			AII
YLR272c-S2	<del>                                     </del>										CCG	CCT
ILRZ/ZC-5Z	TTA										EQ ID	
	172)	900	All	1 00	C CA	.C II	10 10	JO 21	10 1	<u> (5)</u>	EQ ID	
YLR275w-S1	CCG	TTT	TAT	CAT	GTC	GTA	TGT	TTG	ATC	TTA	ACC	ATT
	TTT	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	O NO:	3)	
YLR275w-S2	CAA	CGA	TAA	CTG	AAT	CAC	CTC	TTA	AGA	ATA	GTT	TAC
	TTA	<b>T</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (5	SEQ II	NO:	4)
YLR276c-S1	CTT	CAA	CGG	GTC	TAC	TTT	ACC	ATT	CTT	TGG	CTT	ACT
	GAC	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	ONO:	5)	
YLR276c-S2	AGC	TAT	GAG	AAA	AAG	TCT	GTG	GAA	GGC	GCT	TAT	ATT
	GAC	<b>G</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u>	SEQ II	NO:	6)
YLR317w-S1	CTG	CCA	TCT	TCT	GCC	ACC	ACT	TTG	TCC	TTC	TTT	CTT
	GAT	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	ONO:	7)	
YLR317w-S2	GAA	GTA	AAC	TAA	CTA	GTA	AAG	TAG	GCT	AAT	TCG	AAA
	CGA	TGC	ATA	A GG	C CA	C TA	G TO	GG A	тс т	G (S	EQ ID	NO:
	176)											
YLR359w-S1	GGC	TAT	TGC	TGA	GAA	GGA	ATT	GGG	CTT	AAC	TGT	TGT
	TAC	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	O NO:	8)	
YLR359w-S2	AAC	TTG	ACT	TGT	TCA	TCG	TTT	AGG	TAC	TTT	TGG	AAA
	GGT	<b>T</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u> 9	SEQ II	NO:	9)
YLR373c-S1	ACA	CAC	AGG	TAC	AGA	GTG	CTG	AAA	GAG	GAT	TGG	TGT
	TGC	<b>C</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	NO:	10)	

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YLR373c-S2	CAA	ACA	GAC	TTT	GTT	CCT	TTG	TAT	GTC	CTA	TGG	AAG
	ATA	<b>C</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (S	SEQ ID	NO:	11)
YLR424w-S1	GAC	ATG	ACA	TAC	ACT	AAT	GAT	GCC	TTG	AAA	ACT	AGT
	AGC	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ II	NO:	12)	
YLR424w-S2	ATA	GGT	ACT	TTC	TAG	AGG	TCA	AGG	GCC	CAT	AAA	TAA
	ATT	<b>G</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u> 9	SEQ ID	NO:	13)

B6

Gene deletion	s on	chro	moso	m 1:	2							
Name	Seq	ience	<b>∍</b> 5'-	-3'					-			
YLR437c-S1	ATT	GTG	CAA	GTC	TGT	TAA	AGT	CTT	CTC	TTG	GAT	CCA
	TTA	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ I	D NO:	14)	
YLR437c-S2	CAT	CAC	ACA	CTA	ATA	CAG	GAA	CAA	ACA	AGA	CTT	AAT
	GGA	<b>C</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (:	SEQ II	NO:	15)
YLR440c-S1	TTG	CCA	AGA	AAA	TTG	CAG	TAA	AAA	TGT	TGG	AAG	AGC
	AAC	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ I	D NO:	16)	
YLR440c-S2	GCT	CCA	ATT	CTA	GTG	TGC	TCC	ATT	GCG	ATG	TAA	CAA
	TTT	<b>C</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u> :	SEQ II	NO:	<u>17)</u>

Block

Gene deletions	on	chro	nosoi	me 6								
Name	Sequ	nce	5'-	-3 '								
YFL024c-S1	TGA	TGA	ATT	TTT	CTG	GGT	TAT	AGA	AGA	GTT	CTG	TTT
	CGC	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	O NO:	18)	
YFL024c-S2	ACA	CCT	TCA	AAC	GCT	ATA	GAG	ATC	AAT	GAC	GGT	TCG
	CAT	<b>A</b> GC	ATA	GGC	CAC	TAG	TGG	ATC_	TG <u>(</u>	SEQ II	NO:	20)
YFR003c-S1	TGT	GGA	AGA	GGT	TCC	CGC	AGT	TTT	GCA	GCT	TCG	AGC
	AAC	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	O NO:	21)	
YFR003c-S2	ATC	TTC	TTT	GTC	TAC	GTT	CGT	TAA	AGT	CAA	GAT	CCT
	TCT	<b>C</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u>	SEQ II	NO:	22)
YFR027w-S1	AAT	GAA	AGC	TAG	GAA	ATC	GCA	GAG	AAA	AGC	GGG	CAG
	TAA	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ II	D NO:	23)	
YFR027w-S2	AAT	TTG	GTT	GCG	ATA	CCC	AAC	TTC	CTT	GCT	GTC	CTG
	CAC	<b>A</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (	SEQ II	NO:	24)
YFR042w-S1	AGT	TTG	CAC	CAA	TGG	CAA	TAT	GCC	TGT	GAT	AAA	GAT
	AAG	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ I	D NO:	<u>25)</u>	
YFR042w-S2	CAT	GGA	AGT	TAT	TTG	GTT	GCT	TAG	ATT	CCA	CGG	GTT
	CAA	<b>A</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u>	SEQ II	NO:	26)

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Gene deletions	on	chro	moso	me 9								
Name	S qu	ience	∍ 5'-	-3'		-						
YIL109c-S1	TGT	CTC	ATC	ACA	AGA	AAC	GTG	ттт	ACC	CAC	AAG	CTC
	AGC	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ I	D NO:	27)	
YIL109c-S2	TCA	TGA	TTT	GTA	AGA	ATT	CTC	TGT	AAC	TTT	CGT	TAT
	TCA	<b>A</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(:</u>	SEQ II	NO:	29)
YIL091c-S1	AGT	GAC	AGT	TCT	GTG	AGG	GAA	AAG	AAT	GAT	AAT	TTC
	CGT	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ I	D NO:	30)	
YIL091c-S2	CAT	TGT	AAA	ATT	CAG	GAT	TGT	TTG	GAG	GCT	TAT	AAA
	AAA	<b>C</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (	SEQ II	NO:	31)
YIL083c-S1	ACC	TCT	ACC	CGT	GCT	CAA	CAG	ACC	TCA	AAT	TCA	TAC
	GTC	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ I	D NO:	32)	
YIL083c-S2	CGA	TGA	CTT	CTG	GGA	TTA	TCA	TCT	CTT	CAA	TGA	TAT
	GGT				CAC	TAG	TGG	ATC	TG <u>(</u> :	SEQ II	NO:	33)
YIL019w-S1	TTC	AAA	GAA	AAG	CTT	TTG	AAA	GTC	AGT	TCG	GAT	CTT
	TAG	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ I	D NO:	34)	
YIL019w-S2	TAG	CGA	CGG	GAT	TTC	TTT	TTG	CCA	TTA	AAT	TTA	CCA
	CTC	<b>C</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(</u> :	SEQ II	NO:	<u>35)</u>
YIL104c-S1	TCC	CCT	TAC	TAT	TTA	AGA	TTA	AGA	TTT	CCT	CAC	GAA
	TTA	<b>A</b> CA	GCT						SEQ I			
YIL104c-S2	CCT	GAT	ACC	TGT					CTT			
	GCT	<b>G</b> GC	ATA						TG <u>(</u> :			
YIR010w-S1	ATG	AGT	CTG	_					GTC			ACG
	CCG	<b>C</b> CA		GAA	GCT		TAC		SEQ I			
YIR010w-S2	ACC	TTA		CAT		CTT				ACT		TCT
	CAA	<b>C</b> GC	ATA		CAC		TGG			SEQ II		<del></del>
YIR015w-S1	GCG	AAC	CAG			TTC			TTA		TAC	CTC
	TAC		GCT	GAA				<u>-</u>	SEQ I			
YIR015w-S2	ATT		AGT	TTT					AAC			
	CAT	TGC	ATA	GGC	CAC	TAG	TGG	ATC	TG (	SEQ II	NO:	42)

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Gene deletions	on (	chro	nosor	n 10	5							
Name	Sequ	ence	5'-	-3'								
YPL233w-S1	ATG								CTG			ACT
	GTT	<b>G</b> CA							SEQ II			
YPL233w-S2	CAA	TCC	TCC				TCC			CGC		ACC
	TTT			GGC			TGG			SEQ II		
YPL146c-S1	TCC	AAC	TAA						ATC			CAA
	ACA	<b>G</b> CA		GAA			TAC		SEQ II			
YPL146c-S2	TTT		GTC	CTT	ATG				TTC			
WDT 106 G1	CTG	-	ATA		CAC				TG (S			
YPL126w-S1	ATG		CAA			GGT			CAG SEQ II			CIG
YPL126w-S2	TAT	GCA GTT	AAT	ACT	TTC				TCA			CTA
1PL126W-52	TCC	AGC		GGC		<del>-</del>	TGG		TG (			
YPL093w-S1	CAA	GAT	TAC		LAA		AG(		CTA		GCG	
II DO JA-DI	AGT			_					SEQ II		49)	
YPL093w-S2	CGG			GTC				<del>_</del>	CGC		<del></del>	СТС
1111095# 62	AAT		_				TGG		TG (:			
YPL063w-S1	TTC		CAT				ттт		CAA			
	GTT	TCA		GAA			TAC		SEO I	D NO:	52)	
YPL063w-S2	GGA	TTC	AGC	AAT	CTT				СТТ	TTC		TTC
	AAA	<b>T</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (:	SEQ II	NO:	53)
YPL024w-S1	ATG	TCT	TTT	TCA	TCT				CAG	GAT	ATC	ACA
	GAT	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC (	SEQ I	D NO:	54)	
YPL024w-S2	ACT	TGT	GAG	TCC	TTC	AAT	ATG	AAA	ACG	CCC	CTA	TTG
	AAC	<b>A</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG <u>(:</u>	SEQ II	NO:	55)
YPL020c-S1	TCA	GTT	GAA	GTA	GAT	AAG	CAC	CGG	AAC	ACA	CTA	CAG
	TAT	<b>C</b> CA	GCT	GAA	GCT	TCG	TAC	GC <u>(</u>	SEQ I	D NO:	56)	
YPL020c-S2	TCG	GTT	AAA			TGG	GCA	ATA	AAT	CTT	CTC	ATC
	CTA	<b>A</b> GC		GGC			TGG			SEQ II		
YPL012w-S1	ATG	GAT	CAA		AAA					TTA	GAG	CTG
		<b>G</b> CA					TAC		SEQ I		58)	
YPL012w-S2	ATT		ACT		GAC				TAT			CAA
TEDT 0.07 - 64	<del></del>								TG (:			
YPL007c-S1		_	_						TCT SEQ I			GCG
YPL007c-S2	<del>                                     </del>								ATT			ጥጥር
IPL00/C-52									TG (:			
YPR048w-S1	ATG								ATC			
TEVOSOM-DT									SEQ I			GGA
YPR048w-S2	AAT								TCC			AGT
ZAILUMUW DZ									TG (:			
YPR072w-S1	TGT					-			ATA			
	TTT			_					SEQ I			
YPR072w-S2	AGA								CTT			TTC
									TG (:			
	1											_ <del></del> _

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Name	Seq	uence	3 5'-	-3'								
YPR082c-S1	CTT	CGA	TTG	CTG	AAA	GAG	TAA	GGA	ACT	TTG	CAG	TT
	TTT	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	<u>67)</u>	_
YPR082c-S2	CAA	TAA	AGT	TCA	ACT	TGT	TGT	TGI	TCC	CTG	TAC	CAA
	AAT	<b>C</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (	SEQ II	NO:	68)
YPR085c-S1	CTG	TAC	ATT	CTT	TCG	AAA	GAC	TCC	ATG	CTG	CGA	ATT
	TTT	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	69)	
YPR085c-S2	TCC	CAC	TTT	ATA	GTT	ATG	GGA	ттт	CGA	GCT	GGA	TTC
	GGT	<b>A</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (	SEQ II	NO:	70)
YPR105c-S1	AGC	TCG	ATC	ATC	GAG	GGC	CAA	тто	TCT	AAA	AAT	CTA
	GCA	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	72)	
YPR105c-S2	CTG	TGT	TCT	ATC	AAT	CTT	CAT	ATI	TCT	AGC	TTT	LAA
	TCT	<b>T</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (	SEQ II	NO:	73)
YPR112c-S1	CAT	TGT	CAA	GGG	TTT	GCC	CGT	CTA	тст	AAC	AGA	TGA
	TAA	<b>T</b> CA	GCT	GAA	GCT	TCG	TAC	GC	(SEQ I	D NO:	74)	
YPR112c-S2	GAA	ACC	TTC	GTT	TTC	TTC	ATC	ATC	CAC	ATC	CAG	TTT
	CTT	<b>T</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (	SEQ II	NO:	76)
YPR137w-S1	ATG	TCA	GAT	GTT	ACC	CAA	CAG	AAA	AAG	AGG	AAA	AGA
	TCC	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC	(SEQ I	D NO:	75)	
YPR137w-S2	AAA	AGC	CTG	TTT	GGT	CAA	TGA	CAC	CTG	AAT	ATA	TAC
	CAT	TGC	ATA	GG(	C CA	C TA	G TO	GG A	TC I	G (SI	EQ ID	NO:
	180)											
YPR143w-S1	ATG	GGC	TCC	AAG	CAC	AGA	GTA	GAC	ACT	AAG	GAT	AAG
	AAA	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC _	(SEQ I	D NO:	77)	
YPR143w-S2	TTC	ATT	GTC	GCT	TCC	TGC	GGC	AGC	TTT	AAC	TAA	ATC
	CAA	<b>A</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (	SEQ II	NO:	78)
YPR144c-S1	TTC	CAG	AAA	ATG	TTA	CTC	AAT	TGG	AAG	AAG	ATG	AGA
	CAG	<b>A</b> CA	GCT	GAA	GCT	TCG	TAC	GC _	SEQ I	D NO:	79)	
YPR144c-S2	CCA	TGC	TAC	CCC	AGG	CAA	GTA	GAC	GTT	ACC	TTG	GGA
	TGA	<b>C</b> GC	ATA	GGC	CAC	TAG	TGG	ATC	TG (	SEQ II	NO:	80)
YPR169w-S1	TTT	TAC	ATC	CTG	AAC	TGC	CCA	TTA	TAA	TAA	CTG	GCI
	TTG	<b>G</b> CA	GCT	GAA	GCT	TCG	TAC	GC	SEQ I	D NO:	82)	
YPR169w-S2	СТТ	CTT	GAT	CCC	ATG	CTC	ATA	CAG	GTC	СТТ	TTT	ттт
	GTT	<b>G</b> GC	ΔΤΔ	GGC	CAC	TAG	TGG	ΔТС	TG (	SEO II	NO.	83)

13/